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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/892,635A

DATE: 06/16/2003 P.6

TIME: 15:27:32

Input Set : A:\031998-007.ST25.txt

Output Set: N:\CRF4\06162003\I892635A.raw

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4 <110> APPLICANT: May, Gregory D.
5   Clendennen, Stephanie K.
6   Mason, Hugh S.
7   Lim, Miguel A. Gomez
8   Arntzen, Charles J.
10 <120> TITLE OF INVENTION: DNA Regulatory Elements Associated with Fruit Development
12 <130> FILE REFERENCE: 031998-007
14 <140> CURRENT APPLICATION NUMBER: US 09/892,635A
15 <141> CURRENT FILING DATE: 2001-06-28
17 <150> PRIOR APPLICATION NUMBER: US 09/160,351
18 <151> PRIOR FILING DATE: 1998-09-25
20 <150> PRIOR APPLICATION NUMBER: US 60/060,062
21 <151> PRIOR FILING DATE: 1997-09-25
23 <160> NUMBER OF SEQ ID NOS: 45
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 1186
29 <212> TYPE: DNA
30 <213> ORGANISM: Musa acuminata
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (55)...(1026)
36 <400> SEQUENCE: 1
37 tttggttgtag cctaacagag agagagagac agaccgatag cctcctcatt cact atg      57
38                                     Met
39                                     1
41 gcg atc cga tcg cca gct tcg ctg ctg tta ttt gcg ttc ctg atg ctt      105
42 Ala Ile Arg Ser Pro Ala Ser Leu Leu Leu Phe Ala Phe Leu Met Leu
43           5                      10                      15
45 gcg ctc acg gga aga ctg cag gcc cgg cgc agc tca tgc att ggc gtc      153
46 Ala Leu Thr Gly Arg Leu Gln Ala Arg Arg Ser Ser Cys Ile Gly Val
47           20                      25                      30
49 tac tgg gga caa aac acc gac gag gga agc tta gca gat gct tgt gcc      201
50 Tyr Trp Gly Gln Asn Thr Asp Glu Gly Ser Leu Ala Asp Ala Cys Ala
51           35                      40                      45
53 aca ggc aac tac gaa tac gtg aac atc gcc acc ctt ttc aag ttt ggc      249
54 Thr Gly Asn Tyr Glu Tyr Val Asn Ile Ala Thr Leu Phe Lys Phe Gly
55           50                      55                      60                      65
57 atg ggc caa act cca gag atc aac ctc gcc ggc cac tgt gac cct cgg      297
58 Met Gly Gln Thr Pro Glu Ile Asn Leu Ala Gly His Cys Asp Pro Arg
59           70                      75                      80
61 aac aac ggc tgc gcg cgc ttg agc agc gaa atc cag tcc tgc cag gag      345
62 Asn Asn Gly Cys Ala Arg Leu Ser Ser Glu Ile Gln Ser Cys Gln Glu

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63          85          90          95
65 cgt ggc gtc aag gtg atg ctc tcc atc gga ggt ggc ggg tct tat ggc 393
66 Arg Gly Val Lys Val Met Leu Ser Ile Gly Gly Gly Gly Ser Tyr Gly
67          100          105          110
69 ctg agt tcc acc gaa gac gcc aag gac gta gcg tca tac ctc tgg cac 441
70 Leu Ser Ser Thr Glu Asp Ala Lys Asp Val Ala Ser Tyr Leu Trp His
71          115          120          125
73 agt ttc ttg ggt ggt tct gct gct cgc tac tcg aga ccc ctc ggg gat 489
74 Ser Phe Leu Gly Gly Ser Ala Ala Arg Tyr Ser Arg Pro Leu Gly Asp
75 130          135          140          145
77 gcg gtt ctg gat ggc ata gac ttc aac atc gcc gga ggg agc aca gaa 537
78 Ala Val Leu Asp Gly Ile Asp Phe Asn Ile Ala Gly Gly Ser Thr Glu
79          150          155          160
81 cac tat gat gaa ctt gcc gct ttc ctc aag gcc tac aac gag cag gag 585
82 His Tyr Asp Glu Leu Ala Ala Phe Leu Lys Ala Tyr Asn Glu Gln Glu
83          165          170          175
85 gcc gga acg aag aaa gtt cac ttg agt gct cgt ccg cag tgt cct ttc 633
86 Ala Gly Thr Lys Lys Val His Leu Ser Ala Arg Pro Gln Cys Pro Phe
87          180          185          190
89 ccg gat tac tgg ctt ggc aac gca ctc aga aca gat ctc ttc gac ttc 681
90 Pro Asp Tyr Trp Leu Gly Asn Ala Leu Arg Thr Asp Leu Phe Asp Phe
91          195          200          205
93 gtg tgg gtg cag ttc ttc aac aac cct tcg tgc cat ttc tcc cag aac 729
94 Val Trp Val Gln Phe Phe Asn Asn Pro Ser Cys His Phe Ser Gln Asn
95 210          215          220          225
97 gct atc aat ctt gca aat gcg ttc aac aat tgg gtc atg tcc atc cct 777
98 Ala Ile Asn Leu Ala Asn Ala Phe Asn Asn Trp Val Met Ser Ile Pro
99          230          235          240
101 gcg caa aag ctg ttc ctt ggg ctt cct gct gct cct gag gct gct cca 825
102 Ala Gln Lys Leu Phe Leu Gly Leu Pro Ala Ala Pro Glu Ala Ala Pro
103          245          250          255
105 act ggt ggc tac att cca ccc cat gat ctc ata tct aaa gtt ctt ccg 873
106 Thr Gly Gly Tyr Ile Pro Pro His Asp Leu Ile Ser Lys Val Leu Pro
107          260          265          270
109 atc cta aag gat tcc gac aag tac gca gga atc atg ctg tgg act aga 921
110 Ile Leu Lys Asp Ser Asp Lys Tyr Ala Gly Ile Met Leu Trp Thr Arg
111          275          280          285
113 tac cac gac aga aac tcc ggc tac agt tct caa gtc aag tcc cac gtg 969
114 Tyr His Asp Arg Asn Ser Gly Tyr Ser Ser Gln Val Lys Ser His Val
115 290          295          300          305
117 tgt cca gcg cgt cgg ttc tcc aac atc tta tct atg ccg gtg aag tct 1017
118 Cys Pro Ala Arg Arg Phe Ser Asn Ile Leu Ser Met Pro Val Lys Ser
119          310          315          320
121 tcc aag taa acctgaacgg cgtagatgat cggtggtcga aaactccgat 1066
122 Ser Lys *
125 catcatgggt ccccatccgt atccgtgcgt tgctacgtta tgggtgtttcc cttgtatggt 1126
126 ggtctttttca ataataaat aaggggttag ttttacgttt ccaaaaaaaaa aaaaaaaaaa 1186
129 <210> SEQ ID NO: 2
130 <211> LENGTH: 323

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Input Set : A:\031998-007.ST25.txt

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131 <212> TYPE: PRT
132 <213> ORGANISM: Musa acuminata
134 <400> SEQUENCE: 2
135 Met Ala Ile Arg Ser Pro Ala Ser Leu Leu Leu Phe Ala Phe Leu Met
136 1 5 10 15
137 Leu Ala Leu Thr Gly Arg Leu Gln Ala Arg Arg Ser Ser Cys Ile Gly
138 20 25 30
139 Val Tyr Trp Gly Gln Asn Thr Asp Glu Gly Ser Leu Ala Asp Ala Cys
140 35 40 45
141 Ala Thr Gly Asn Tyr Glu Tyr Val Asn Ile Ala Thr Leu Phe Lys Phe
142 50 55 60
143 Gly Met Gly Gln Thr Pro Glu Ile Asn Leu Ala Gly His Cys Asp Pro
144 65 70 75 80
145 Arg Asn Asn Gly Cys Ala Arg Leu Ser Ser Glu Ile Gln Ser Cys Gln
146 85 90 95
147 Glu Arg Gly Val Lys Val Met Leu Ser Ile Gly Gly Gly Gly Ser Tyr
148 100 105 110
149 Gly Leu Ser Ser Thr Glu Asp Ala Lys Asp Val Ala Ser Tyr Leu Trp
150 115 120 125
151 His Ser Phe Leu Gly Gly Ser Ala Ala Arg Tyr Ser Arg Pro Leu Gly
152 130 135 140
153 Asp Ala Val Leu Asp Gly Ile Asp Phe Asn Ile Ala Gly Gly Ser Thr
154 145 150 155 160
155 Glu His Tyr Asp Glu Leu Ala Ala Phe Leu Lys Ala Tyr Asn Glu Gln
156 165 170 175
157 Glu Ala Gly Thr Lys Lys Val His Leu Ser Ala Arg Pro Gln Cys Pro
158 180 185 190
159 Phe Pro Asp Tyr Trp Leu Gly Asn Ala Leu Arg Thr Asp Leu Phe Asp
160 195 200 205
161 Phe Val Trp Val Gln Phe Phe Asn Asn Pro Ser Cys His Phe Ser Gln
162 210 215 220
163 Asn Ala Ile Asn Leu Ala Asn Ala Phe Asn Asn Trp Val Met Ser Ile
164 225 230 235 240
165 Pro Ala Gln Lys Leu Phe Leu Gly Leu Pro Ala Ala Pro Glu Ala Ala
166 245 250 255
167 Pro Thr Gly Gly Tyr Ile Pro Pro His Asp Leu Ile Ser Lys Val Leu
168 260 265 270
169 Pro Ile Leu Lys Asp Ser Asp Lys Tyr Ala Gly Ile Met Leu Trp Thr
170 275 280 285
171 Arg Tyr His Asp Arg Asn Ser Gly Tyr Ser Ser Gln Val Lys Ser His
172 290 295 300
173 Val Cys Pro Ala Arg Arg Phe Ser Asn Ile Leu Ser Met Pro Val Lys
174 305 310 315 320
175 Ser Ser Lys
179 <210> SEQ ID NO: 3
180 <211> LENGTH: 90
181 <212> TYPE: PRT
182 <213> ORGANISM: Musa acuminata
184 <400> SEQUENCE: 3

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Output Set: N:\CRF4\06162003\I892635A.raw

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185 Met Ala Ile Arg Ser Pro Ala Ser Leu Leu Leu Phe Ala Phe Leu Met
186 1 5 10 15
187 Leu Ala Leu Thr Gly Arg Leu Gln Ala Arg Arg Ser Ser Cys Ile Gly
188 20 25 30
189 Val Tyr Trp Gly Gln Asn Thr Asp Glu Gly Ser Leu Ser Asp Lys Tyr
190 35 40 45
191 Ala Gly Ile Met Leu Trp Thr Arg Tyr His Asp Arg Asn Ser Gly Tyr
192 50 55 60
193 Ser Ser Gln Val Lys Ser His Val Cys Pro Ala Arg Arg Phe Ser Asn
194 65 70 75 80
195 Ile Leu Ser Met Pro Val Lys Ser Ser Lys
196 85 90
199 <210> SEQ ID NO: 4
200 <211> LENGTH: 67
201 <212> TYPE: PRT
202 <213> ORGANISM: Musa acuminata
204 <400> SEQUENCE: 4
205 Met Glu Lys Cys Phe Asn Ile Ile Pro Ser Leu Leu Leu Ile Ser Leu
206 1 5 10 15
207 Leu Ile Lys Ser Ser Asn Ala Ala Gly Ile Ala Val Tyr Trp Gly Gln
208 20 25 30
209 Asn Gly Asn Glu Gly Ser Leu Ser Pro Lys Tyr Gly Gly Val Met Ile
210 35 40 45
211 Trp Asp Arg Phe Asn Asp Ala Gln Ser Gly Tyr Ser Asn Ala Ile Lys
212 50 55 60
213 Gly Ser Val
214 65
217 <210> SEQ ID NO: 5
218 <211> LENGTH: 69
219 <212> TYPE: PRT
220 <213> ORGANISM: Musa acuminata
222 <400> SEQUENCE: 5
223 Met Ala Arg Thr Pro Gln Ser Thr Pro Leu Leu Ile Ser Leu Ser Val
224 1 5 10 15
225 Leu Ala Leu Ile Lys Thr Ser Tyr Ala Gly Gly Ile Ala Ile Tyr Trp
226 20 25 30
227 Gly Gln Asn Gly Asn Glu Gly Thr Leu Ser Pro Lys Tyr Gly Gly Val
228 35 40 45
229 Met Ile Trp Ser Lys Phe Tyr Asp Asp Gln Ser Gly Tyr Ser Asn Ser
230 50 55 60
231 Ile Lys Gly Ser Val
232 65
235 <210> SEQ ID NO: 6
236 <211> LENGTH: 73
237 <212> TYPE: PRT
238 <213> ORGANISM: Musa acuminata
240 <400> SEQUENCE: 6
241 Met Thr Asn Met Thr Leu Arg Lys His Val Ile Tyr Pro Leu Leu Phe
242 1 5 10 15

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Input Set : A:\031998-007.ST25.txt

Output Set: N:\CRF4\06162003\I892635A.raw

243 Ile Ser Cys Ser Leu Ser Lys Pro Ser Asp Ala Ser Arg Gly Gly Ile
 244 20 25 30
 245 Ala Ile Tyr Trp Gly Gln Asn Gly Asn Glu Gly Asn Leu Ser Arg Lys
 246 35 40 45
 247 Tyr Gly Gly Val Met Ile Trp Ser Lys Phe Trp Asp Asp Lys Asn Gly
 248 50 55 60
 249 Tyr Ser Asn Ser Ile Leu Ala Ser Val
 250 65 70

253 <210> SEQ ID NO: 7

254 <211> LENGTH: 64

255 <212> TYPE: PRT

256 <213> ORGANISM: Musa acuminata

258 <400> SEQUENCE: 7

259 Met Ile Lys Tyr Ser Pro Leu Leu Thr Ala Ser Val Ser Phe Leu Lys
 260 1 5 10 15
 261 Ala Leu Lys Leu Glu Ala Gly Asp Ile Val Ile Tyr Trp Gly Gln Asn
 262 20 25 30
 263 Gly Asn Glu Gly Asn Leu Ser Pro Lys Tyr Gly Gly Val Met Ile Trp
 264 35 40 45
 265 Ser Lys Phe Tyr Asp Asn Gly Tyr Ser Asn Ala Ile Leu Ala Asn Val
 266 50 55 60

269 <210> SEQ ID NO: 8

270 <211> LENGTH: 67

271 <212> TYPE: PRT

272 <213> ORGANISM: Musa acuminata

274 <400> SEQUENCE: 8

275 Met Ala Ala Lys Ile Val Ser Val Leu Phe Leu Ile Ser Ser Leu Ile
 276 1 5 10 15
 277 Phe Ala Ser Phe Glu Ser Ser His Gly Gly Gln Ile Val Ile Tyr Trp
 278 20 25 30
 279 Gly Gln Asn Gly Asn Glu Gly Asn Leu Ser Ala Lys Tyr Gly Gly Val
 280 35 40 45
 281 Met Ile Trp Ser Lys Ala Tyr Asp Asn Gly Tyr Ser Asn Ala Ile Leu
 282 50 55 60
 283 Ala Ser Val
 284 65

287 <210> SEQ ID NO: 9

288 <211> LENGTH: 496

289 <212> TYPE: DNA

290 <213> ORGANISM: Musa acuminata

292 <220> FEATURE:

293 <221> NAME/KEY: misc_feature

294 <222> LOCATION: 163, 387, 471

295 <223> OTHER INFORMATION: n = A,T,C or G

297 <400> SEQUENCE: 9

298 ggcacgagta catcctctgc ttcttcgagc cttttcgcct tccttcctcg tctaaccatg 60
 299 tcgacctgcg gcaactgcga ctgcgttgac aagagccagt gcgtgaagaa gggaaacagc 120
 W--> 300 tacggatatcg atattgttga gaccgagaag agctacgtcg acnaggtgat cgttgccgca 180
 301 gaagctgccg agcatgacgg caagtgcgaag tgcggcgccg cctgcgcctg caccgactgc 240

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/892,635ADATE: 06/16/2003
TIME: 15:27:33Input Set : A:\031998-007.ST25.txt
Output Set: N:\CRF4\06162003\I892635A.raw**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; N Pos. 163,387,471
Seq#:22; N Pos. 82,601,628,640,655,692,725,774,793,806,813,854,867,870,876
Seq#:22; N Pos. 882,890,919,946,959,965,995,999,1002,1028,1043,1054,1075
Seq#:22; N Pos. 1093,1515,2166,2216,2265,2345,2533,2870,2917,3077,3337,3356
Seq#:22; N Pos. 3618,3627,3754,3810,3819,3884,3893,4494,4503,4524,4533,4568
Seq#:22; N Pos. 4574,4597,4654,4724,4741,4759,4852,5027,5253,5546,5565,5567
Seq#:22; N Pos. 5575,5578,5618,5619,5650,5669,5672,5677,5683,5694,5704,5708
Seq#:22; N Pos. 5732,5741,5754,5758,5772,5778,5780,5784,5788,5802,5804,5808
Seq#:22; N Pos. 5813,5820,5824,5832,5834,5836,5854,5858,5863,5872,5875,5889
Seq#:22; N Pos. 5915,5922,5950,5990,6006,6011,6344,6401,6416,6596,6600,6608
Seq#:22; N Pos. 6612,6712,6748,6753,6756,6762,6830,6844,6847,6863,6910,6965
Seq#:22; N Pos. 6968,7070,7116,7179,7291,7322,7325,7345,7351,7359,7387,7395
Seq#:23; N Pos. 82,601,628,640,655,692,725,774,793,806,813,854,867,870,876
Seq#:23; N Pos. 882,890,919,946,959,965,995,999,1002,1028,1043,1054,1075
Seq#:23; N Pos. 1093,1515,2166,2216,2265,2345,2533,2870,2917,3077,3337,3356
Seq#:23; N Pos. 3618,3627,3754,3810,3819,3884,3893,4494,4503,4524,4533,4568
Seq#:23; N Pos. 4574,4597,4654,4724,4741,4759,4852,5027,5253,5546,5565,5567
Seq#:23; N Pos. 5575,5578,5618,5619,5650,5669,5672,5677,5683,5694,5704,5708
Seq#:23; N Pos. 5732,5741,5754,5758,5772,5778,5780,5784,5788,5802,5804,5808
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Seq#:23; N Pos. 5915,5922,5950,5990,6006,6011,6344,6401,6416,6596,6600,6608
Seq#:23; N Pos. 6612,6712,6748,6753,6756,6762,6830,6844,6847,6863,6910,6965
Seq#:23; N Pos. 6968,7070,7116,7179,7291,7322,7325,7345,7351,7359,7387,7395
Seq#:24; Xaa Pos. 25,164,173,177,181,193,204,220,227,231,233,247,251,254
Seq#:24; Xaa Pos. 259,269,278,282,284,294,296,305,310,314,320,326,458,656
Seq#:24; Xaa Pos. 673,687,713,774,883,899,952,1038,1043,1163,1180,1183,1202
Seq#:24; Xaa Pos. 1204,1397,1400,1412,1414,1422,1441,1462,1468,1474,1504
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Seq#:24; Xaa Pos. 2221,2236,2257,2293,2303,2304,2310,2312,2323
Seq#:25; Xaa Pos. 27,160,169,173,190,200,216,222,227,229,243,247,248,250
Seq#:25; Xaa Pos. 252,255,273,278,280,290,291,292,301,306,459,654,669,685
Seq#:25; Xaa Pos. 710,767,878,893,944,1029,1104,1107,1162,1164,1182,1185
Seq#:25; Xaa Pos. 1372,1375,1379,1382,1393,1395,1417,1440,1445,1450,1475
Seq#:25; Xaa Pos. 1533,1666,1672,1673,1676,1690,1706,1707,1714,1719,1727
Seq#:25; Xaa Pos. 1730,1734,1740,1742,1743,1744,1750,1751,1752,1754,1756
Seq#:25; Xaa Pos. 1757,1760,1761,1769,1770,1773,1779,1788,1790,1799,1813
Seq#:25; Xaa Pos. 1818,1820,1930,1949,1954,2011,2012,2015,2016,2058,2060
Seq#:25; Xaa Pos. 2061,2063,2083,2087,2088,2092,2106,2122,2123,2156,2171
Seq#:25; Xaa Pos. 2191,2224,2235,2236,2247,2258
Seq#:26; Xaa Pos. 23,196,205,209,214,226,254,260,264,267,284,285,287,289

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : **A:\031998-007.ST25.txt**
Output Set: **N:\CRF4\06162003\I892635A.raw**

Seq#:26; Xaa Pos. 301,309,313,315,327,328,345,352,358,494,697,713,728,813
Seq#:26; Xaa Pos. 939,991,1076,1081,1165,1167,1208,1226,1229,1250,1253,1442
Seq#:26; Xaa Pos. 1445,1452,1455,1465,1474,1492,1511,1517,1523,1549,1604
Seq#:26; Xaa Pos. 1676,1770,1777,1780,1781,1794,1795,1805,1812,1814,1816
Seq#:26; Xaa Pos. 1820,1823,1832,1835,1840,1841,1846,1847,1849,1850,1855
Seq#:26; Xaa Pos. 1857,1861,1862,1864,1865,1871,1872,1874,1877,1878,1883
Seq#:26; Xaa Pos. 1894,1903,1916,1922,1923,2034,2053,2058,2114,2118,2144

VERIFICATION SUMMARY

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Input Set : A:\031998-007.ST25.txt

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L:300 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:120
M:341 Repeated in SeqNo=9
L:1050 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:1057 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:1062 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:1064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:60
M:341 Repeated in SeqNo=22
L:1210 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:1217 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:1222 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:1224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:60
M:341 Repeated in SeqNo=23
L:1370 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:1377 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:1380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:16
M:341 Repeated in SeqNo=24
L:1692 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:1698 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:16
M:341 Repeated in SeqNo=25
L:2005 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:2012 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:2015 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:16
M:341 Repeated in SeqNo=26
L:2336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:840
M:341 Repeated in SeqNo=27
L:2431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:840
M:341 Repeated in SeqNo=28
L:2658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:1168
M:341 Repeated in SeqNo=29
L:2739 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:144
M:341 Repeated in SeqNo=30
L:2952 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:160
M:341 Repeated in SeqNo=31
L:3169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:1680
L:3221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:1680
L:3313 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:544
L:3420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:528
L:3531 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:544
L:3600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:1680
M:341 Repeated in SeqNo=37
L:3644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:1680
M:341 Repeated in SeqNo=38
L:3728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:544
M:341 Repeated in SeqNo=39
L:3815 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:528
M:341 Repeated in SeqNo=40
L:3936 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:840

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Input Set : A:\031998-007.ST25.txt

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L:3984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:840